

AMENDMENTS

Listing of claims

The listing of claims will replace all prior versions, and listings of claims in the application.

- 1-9. (cancelled)
10. (currently amended) The method of claim [[9]] 17, wherein said mRNA is amplified by an RNA polymerase reaction.
11. (currently amended) The method of claim [[9]] 17, wherein said mRNA is amplified by reverse transcriptase polymerase chain reaction or the ligase chain reaction.
12. (currently amended) The method of claim [[9]] 17, wherein said detecting is by RNA fingerprinting, branched DNA or a nuclease protection assay.
- 13-14. (cancelled)
15. (currently amended) The method of claim [[1]] 17, wherein the metastatic cancer is metastatic prostate cancer.
16. (currently amended) The method of claim [[1,]] 17, wherein the metastatic cancer is metastatic breast cancer.
17. (currently amended) A method of detecting metastatic cancer, comprising the steps of:
 - (a) detecting the quantity of a disease marker mRNA expressed in human peripheral blood; and

(b) comparing the quantity of said marker to the quantity expressed in peripheral blood of a normal individual;

wherein ~~The method of claim 9, in which~~ said mRNA comprises one or more of the sequences or the complements of the sequences disclosed herein as SEQ ID NO: 1, SEQ ID NO:2, SEQ ID NO:3, SEQ ID NO:29, SEQ ID NO:34, SEQ ID NO:48 or SEQ ID NO:49 and wherein a difference in quantity of expression is indicative of metastatic cancer.

18. (cancelled)

19. (currently amended) The method of claim [[9]] 17, further comprising the steps of

- a) providing primers that selectively amplify said metastatic cancer marker;
- b) amplifying said nucleic acid with said primers to form nucleic acid amplification products;
- c) detecting said nucleic acid amplification products; and
- d) measuring the amount of said nucleic acid amplification products formed.

20. (original) The method of claim 19 in which said primers are selected to specifically amplify a nucleic acid having a sequence comprising SEQ ID NO:1, SEQ ID NO:2, SEQ ID NO:3, SEQ ID NO:29, SEQ ID NO:34, SEQ ID NO:48 or SEQ ID NO:49.

21. (withdrawn) The method of claim 8, wherein said marker is a polypeptide.

22. (withdrawn) The method of claim 21, wherein said polypeptide is encoded by a nucleic acid sequence comprising the sequence disclosed herein SEQ ID NO:1, SEQ ID NO:2, SEQ ID NO: ID NO:3, SEQ ID NO:29, SEQ ID NO:34, SEQ ID NO:48 or SEQ ID NO:49.
23. (withdrawn) The method of claim 21, wherein said detection is by an antibody immunoreactive with said marker.
24. (withdrawn) The method of claim 21, wherein said polypeptide is encoded by an IL-8 or IL-10 gene.
25. (withdrawn) The method of claim 8, wherein said marker is a product of the IL-8 gene and wherein said comparison is between two alternatively spliced forms of an IL-8 gene product.
26. (withdrawn) The method of claim 24, wherein the quantity of IL-8 polypeptide in peripheral blood is measured using an in vitro bioassay that detects an IL-8 mediated biological process.
- 27-64. (cancelled)
65. (withdrawn) The method of claim 24, wherein the quantity of IL-10 polypeptide in peripheral blood is measured using an in vitro bioassay that detects at least one IL-10 mediated biological process.
- 66-73. (cancelled)